IAP5 Rec'd PCT/PTO 01 SEP 2006

SEQUENCE LISTING

<110> Frohberg, Claus Koetting, Oliver

> Ritte, Gerhard Steup, Martin

<120> Methods for identifying proteins with starch phosphorylating enzymatic activity

- <130> 65084.000020
- <140> Unassigned <141> 2006-09-01
- <150> PCT/EP2005/002454
- <151> 2005-03-04
- <150> EP04090483.1
- <151> 2004-12-15
- <150> EP04090121.7
- <151> 2004-03-29
- <150> EP04090087.0
- <151> 2004-03-05
- <150> US60/549,980 provisional
- <151> 2004-03-05
- <160> 46
- <170> PatentIn version 3.1
- <210> 1
- <211>· 3591
- <212> DNA

<213> Arabidopsis thaliana

<220>

<221> CDS (1)..(3591)<222> <223> <400> 1 atg gag agc att ggc agc cat tgt tgc agc tct cct ttc acc ttc atc 48 Met Glu Ser Ile Gly Ser His Cys Cys Ser Ser Pro Phe Thr Phe Ile 15 act aga aac tca tca tca ctt cct aga ctc gtt aac atc act cac 96 Thr Arg Asn Ser Ser Ser Leu Pro Arg Leu Val Asn Ile Thr His 20 aga gtt aat ctc agc cac caa tct cac cga ctc aga aac tcc aat tct 144 Arg Val Asn Leu Ser His Gln Ser His Arg Leu Arg Asn Ser Asn Ser 40 cgt ctc act tgc act gct act tct tct tcc acc att gag gaa caa cgg 192 Arg Leu Thr Cys Thr Ala Thr Ser Ser Ser Thr Ile Glu Glu Gln Arg 50 55 60 aag aag aaa gat gga tca gga acg aaa gtg agg ttg aat gtg agg tta 240 Lys Lys Lys Asp Gly Ser Gly Thr Lys Val Arg Leu Asn Val Arg Leu 80 gat cat caa gtt aat ttt ggt gac cat gtg gct atg ttt gga tca gct 288 Asp His Gln Val Asn Phe Gly Asp His Val Ala Met Phe Gly Ser Ala aaa gag att ggt tca tgg aaa aag aaa tcg cct ttg aat tgg agt gag 336 Lys Glu Ile Gly Ser Trp Lys Lys Ser Pro Leu Asn Trp Ser Glu 100 aat gga tgg gtt tgt gag ttg gaa ctt gac ggt ggt cag gtt ttg gag 384 Asn Gly Trp Val Cys Glu Leu Glu Leu Asp Gly Gly Gln Val Leu Glu 120 125 tat aag ttt gtc att gtt aag aat gat ggt tca ctt tca tgg gaa tct 432 Tyr Lys Phe Val Ile Val Lys Asn Asp Gly Ser Leu Ser Trp Glu Ser 130 140 ggt gat aat cgt gtc ctt aag gtt cca aat tct ggg aat ttt tct gtt 480 Gly Asp Asn Arg Val Leu Lys Val Pro Asn Ser Gly Asn Phe Ser Val 145 150 155 160 gtt tgt cat tgg gat gct act aga gaa acc ctt gat ttg cct cag gag 528 Val Cys His Trp Asp Ala Thr Arg Glu Thr Leu Asp Leu Pro Gln Glu 165 170 175 gtt ggt aat gat gat gtt ggt gat ggg cat gag agg gat aat 576 Val Gly Asn Asp Asp Asp Val Gly Asp Gly Gly His Glu Arg Asp Asn 180 185 190

Н	at Iis	gat Asp	gtt Val 195	ggt Gly	gat Asp	gat Asp	aga Arg	gta Val 200	gtg Val	gga Gly	agt Ser	gaa Glu	aat Asn 205	ggt Gly	gcg Ala	cag Gln	624	
C L	tt .eu	cag Gln 210	aag Lys	agt Ser	aca Thr	ttg Leu	ggt Gly 215	ggg Gly	caa Gln	tgg Trp	caa Gln	ggt Gly 220	aaa Lys	gat Asp	gcg Ala	tcc Ser	672	
Р	tt he	atg Met	cgt Arg	tct Ser	aat Asn	gat Asp 230	cat His	ggt Gly	aac Asn	aga Arg	gaa Glu 235	gtt Val	ggt Gly	aga Arg	aat Asn	tgg Trp 240	720	
	jat NSP	act Thr	agt Ser	ggt Gly	ctt Leu 245	gaa Glu	ggc Gly	aca Thr	gct Ala	ctt Leu 250	aag Lys	atg Met	gtt Val	gag Glu	ggt Gly 255	gat Asp	768	
C A	gc rg	aac Asn	tct Ser	aag Lys 260	aac Asn	tgg Trp	tgg Trp	aga Arg	aag Lys 265	ctt Leu	gaa Glu	atg Met	gta Val	cgc Arg 270	gag Glu	gtt Val	816	
a	ita He	gtt Val	ggg Gly 275	agt Ser	gtt Val	gag Glu	agg Arg	gag Glu 280	gaa Glu	cga Arg	ttg Leu	aag Lys	gcg Ala 285	ctc Leu	ata Ile	tac Tyr	864	
t S	ct	gca Ala 290	att Ile	tat Tyr	ttg Leu	aag Lys	tgg Trp 295	ata Ile	aac Asn	aca Thr	ggt Gly	cag Gln 300	att Ile	cct Pro	tgt Cys	ttt Phe	912	
G	jaa Slu 805	gat Asp	gga Gly	ggg Gly	cat His	cac His 310	cgt Arg	cca Pro	aac Asn	agg Arg	cat His 315	gcc Ala	gag Glu	att Ile	tcc Ser	aga Arg 320	960	
L	tt .eu	ata Ile	ttc Phe	cgt Arg	gag Glu 325	ttg Leu	gag Glu	cac His	att Ile	tgc Cys 330	agt Ser	aag Lys	aaa Lys	gat Asp	gct Ala 335	act Thr	1008	
P	ca Pro	gag Glu	gaa Glu	gtg Val 340	ctt Leu	gtt Val	gct Ala	cgg Arg	aaa Lys 345	atc Ile	cat His	ccg Pro	tgt Cys	tta Leu 350	cct Pro	tct Ser	1056	
Þ	tc	aaa Lys	gca Ala 355	gag Glu	ttt Phe	act Thr	gca Ala	gct Ala 360	gtc Val	cct Pro	cta Leu	act Thr	cgg Arg 365	att Ile	agg Arg	gac Asp	1104	
a	ita :le	gcc Ala 370	cat His	cgg Arg	aat Asn	gat Asp	att Ile 375	cct Pro	cat His	gat Asp	ctc Leu	aag Lys 380	caa Gln	gaa Glu	atc Ile	aag Lys	1152	
H	at lis 85	acg Thr	ata Ile	caa Gln	aat Asn	aag Lys 390	ctt Leu	cac His	cgg Arg	aat Asn	gct Ala 395	ggt Gly	cca Pro	gaa Glu	gat Asp	cta Leu 400	1200	
a I	itt le	gca Ala	aca Thr	gaa Glu	gca Ala 405	atg Met	ctt Leu	caa Gln	cga Arg	att Ile 410	acc Thr	gag Glu	acc Thr	cca Pro	gga Gly 415	aaa Lys	1248	
t T	at	agt Ser	gga Gly	gac Asp 420	ttt Phe	gtg Val	gag Glu	cag Gln	ttt Phe 425	aaa Lys	ata Ile	ttc Phe	cat His	aat Asn 430	gag Glu	ctt Leu	1296	
a L	aa .ys	gat Asp	ttc Phe 435	ttt Phe	aat Asn	gct Ala	gga Gly	agt Ser 440	ctc Leu	act Thr	gaa Glu	cag Gln	ctt Leu 445	gat Asp	tct Ser	atg Met	1344	

aaa Lys	att Ile 450	tct Ser	atg Met	gat Asp	gat Asp	aga Arg 455	ggt Gly	ctt Leu	tct Ser	gcg Ala	ctc Leu 460	aat Asn	ttg Leu	ttt Phe	ttt Phe	1392
gaa Glu 465	tgt Cys	aaa Lys	aag Lys	cgc Arg	ctt Leu 470	gac Asp	aca Thr	tca Ser	gga Gly	gaa Glu 475	tca Ser	agc Ser	aat Asn	gtt Val	ttg Leu 480	1440
gag Glu	ttg Leu	att Ile	aaa Lys	acc Thr 485	atg Met	cat His	tct Ser	cta Leu	gct Ala 490	tct Ser	tta Leu	aga Arg	gaa Glu	aca Thr 495	att Ile	1488
ata Ile	aag Lys	gaa Glu	ctt Leu 500	aat Asn	agc Ser	ggc Gly	ttg Leu	cga Arg 505	aat Asn	gat Asp	gct Ala	cct Pro	gat Asp 510	act Thr	gcc Ala	1536
att Ile	gca Ala	atg Met 515	cgc Arg	cag Gln	aag Lys	tgg Trp	cgc Arg 520	ctt Leu	tgt Cys	gag Glu	atc Ile	ggc Gly 525	ctc Leu	gag Glu	gac Asp	1584
tac Tyr	ttt Phe 530	ttt Phe	gtt Val	cta Leu	cta Leu	agc Ser 535	aga Arg	ttc Phe	ctc Leu	aat Asn	gct Ala 540	ctt Leu	gaa Glu	act Thr	atg Met	1632
gga Gly 545	gga Gly	gct Ala	gat Asp	caa Gln	ctg Leu 550	gca Ala	aaa Lys	gat Asp	gtg val	gga Gly 555	tca Ser	aga Arg	aac Asn	gtt Val	gcc Ala 560	1680
tca Ser	tgg Trp	aat Asn	gat Asp	cca Pro 565	cta Leu	gat Asp	gct Ala	ttg Leu	gtg Val 570	ttg Leu	ggt Gly	gtt Val	cac His	caa Gln 575	gta Val	1728
ggt Gly	cta Leu	tct Ser	ggt Gly 580	tgg Trp	aag Lys	caa Gln	gaa Glu	gaa Glu 585	tgt Cys	tta Leu	gcc Ala	att Ile	gga Gly 590	aat Asn	gaa Glu	1776
ctc Leu	ctt Leu	gct Ala 595	tgg Trp	cga Arg	gaa Glu	agg Arg	gac Asp 600	cta Leu	ctt Leu	gaa Glu	aaa Lys	gaa Glu 605	ggg Gly	gaa Glu	gag Glu	1824
gat Asp	gga Gly 610	aaa Lys	aca Thr	att Ile	tgg Trp	gcc Ala 615	atg Met	agg Arg	ctg Leu	aaa Lys	gca Ala 620	act Thr	ctt Leu	gat Asp	cga Arg	1872
gca Ala 625	cgc Arg	aga Arg	tta Leu	aca Thr	gca Ala 630	gaa Glu	tat Tyr	tct Ser	gat Asp	ttg Leu 635	ctt Leu	ctt Leu	caa Gln	ata Ile	ttt Phe 640	1920
cct Pro	cct Pro	aat Asn	gtg Val	gag Glu 645	att Ile	tta Leu	gga Gly	aaa Lys	gct Ala 650	cta Leu	gga Gly	att Ile	cca Pro	gag Glu 655	aat Asn	1968
agt Ser	gtc Val	aag Lys	acc Thr 660	tat Tyr	aca Thr	gaa Glu	gca Ala	gag Glu 665	att Ile	cgt Arg	gct Ala	gga Gly	att Ile 670	att Ile	ttc Phe	2016
cag Gln	atc Ile	tca Ser 675	aag Lys	ctc Leu	tgc Cys	act Thr	gtt Val 680	ctt Leu	cta Leu	aaa Lys	gct Ala	gta Val 685	aga Arg	aat Asn	tca Ser	2064
ctt Leu	ggt Gly 690	tct Ser	gag Glu	ggc Gly	tgg Trp	gat Asp 695	gtc Val	gtt Val	gta Val	cct Pro	gga Gly 700	tcg Ser	acg Thr	tct Ser	ggg Gly	2112

	aca Thr 705	tta Leu	gtt Val	cag Gln	gtt Val	gag Glu 710	agc Ser	att Ile	gtt Val	ccg Pro	gga Gly 715	tca Ser	ttg Leu	cca Pro	gca Ala	act Thr 720	2160
1	tct Ser	ggt Gly	ggt Gly	cct Pro	att Ile 725	att Ile	ctc Leu	ttg Leu	gtc Val	aat Asn 730	aaa Lys	gct Ala	gat Asp	ggc Gly	gat Asp 735	gaa Glu	2208
(gag Glu	gta Val	agt Ser	gct Ala 740	gct Ala	aat Asn	ggg Gly	aac Asn	ata Ile 745	gct Ala	gga Gly	gtc Val	atg Met	ctt Leu 750	ctg Leu	cag Gln	2256
(gag Glu	ctg Leu	cct Pro 755	cac His	ttg Leu	tct Ser	cac His	ctt Leu 760	ggc Gly	gtt Val	aga Arg	gcg Ala	cgg Arg 765	cag Gln	gag Glu	aaa Lys	2304
•	att Ile	gtc Val 770	ttt Phe	gtg Val	aca Thr	tgt Cys	gat Asp 775	gat Asp	gat Asp	gac Asp	aag Lys	gtt Val 780	gct Ala	gat Asp	ata Ile	cga Arg	2352
1	cga Arg 785	ctt Leu	gtg Val	gga Gly	aaa Lys	ttt Phe 790	gtg Val	agg Arg	ttg Leu	gaa Glu	gca Ala 795	tct Ser	cca Pro	agt Ser	cat His	gtg Val 800	2400
į	aat Asn	ctg Leu	ata Ile	ctt Leu	tca Ser 805	act Thr	gag Glu	ggt Gly	agg Arg	agt Ser 810	cgc Arg	act Thr	tcc Ser	aaa Lys	tcc Ser 815	agt Ser	2448
						gat Asp											2496
1	aag Lys	aag Lys	agc Ser 835	tta Leu	tct Ser	atc Ile	gat Asp	gat Asp 840	gaa Glu	gaa Glu	tca Ser	aag Lys	cct Pro 845	ggt Gly	tcc Ser	tca Ser	2544
						ctt Leu											2592
						gat Asp 870											2640
(gct Ala	gca Ala	tgt Cys	ggt Gly	ctt Leu 885	ctt Leu	gca Ala	tct Ser	tta Leu	gca Ala 890	gaa Glu	gcc Ala	tct Ser	agt Ser	aaa Lys 895	gtg Val	2688
						gtt Val											2736
,	gtc Val	ata Ile	cct Pro 915	ttt Phe	gga Gly	tcg Ser	atg Met	gaa Glu 920	tta Leu	gct Ala	tta Leu	aag Lys	caa Gln 925	aat Asn	aat Asn	tcg Ser	2784
						tct Ser											2832
(gag Glu 945	ggt Gly	ggt Gly	gag Glu	cta Leu	gac Asp 950	gac Asp	ata Ile	tgt Cys	gac Asp	cag Gln 955	atc Ile	cat His	gaa Glu	gtg Val	atg Met 960	2880

aaa acg ttg caa gtg cct aaa gaa aca atc aac agc ata agc aaa gcg Lys Thr Leu Gln Val Pro Lys Glu Thr Ile Asn Ser Ile Ser Lys Ala 965 970 975	2928
ttt ctc aaa gat gct cgt ctc att gtt cgt tca agt gct aac gtc gag Phe Leu Lys Asp Ala Arg Leu Ile Val Arg Ser Ser Ala Asn Val Glu 980 985 990	2976
gac tta gcc gga atg tca gct gca gga ctc tat gaa tca atc cct aac Asp Leu Ala Gly Met Ser Ala Ala Gly Leu Tyr Glu Ser Ile Pro Asn 995 1000 1005	3024
gtg agt ccc tcg gat cct ttg gtg ttt tca gat tcg gtt tgc caa Val Ser Pro Ser Asp Pro Leu Val Phe Ser Asp Ser Val Cys Gln 1010 1015 1020	3069
gtt tgg gct tct ctc tac aca aga aga gct gtt cta agc cgt aga Val Trp Ala Ser Leu Tyr Thr Arg Arg Ala Val Leu Ser Arg Arg 1025 1030 1035	3114
gct gct ggt gtc tct caa aga gaa gct tca atg gct gtt ctc gtt Ala Ala Gly Val Ser Gln Arg Glu Ala Ser Met Ala Val Leu Val 1040 1045 1050	3159
caa gaa atg ctt tcg ccg gac tta tca ttc gtt ctg cac aca gtg Gln Glu Met Leu Ser Pro Asp Leu Ser Phe Val Leu His Thr Val 1055 1060 1065	3204
agt cca gct gat ccg gac agt aac ctt gtg gaa gcc gag atc gct Ser Pro Ala Asp Pro Asp Ser Asn Leu Val Glu Ala Glu Ile Ala 1070 1075 1080	3249
cct ggt tta ggt gag act tta gct tca gga aca aga gga aca cca Pro Gly Leu Gly Glu Thr Leu Ala Ser Gly Thr Arg Gly Thr Pro 1085 1090 1095	3294
tgg aga ctc gct tcg ggt aag ctc gac ggg att gta caa acc tta Trp Arg Leu Ala Ser Gly Lys Leu Asp Gly Ile Val Gln Thr Leu 1100 1105 1110	3339
gct ttc gca aac ttc agc gaa gag ctt ctt gtg tca gga aca ggt Ala Phe Ala Asn Phe Ser Glu Glu Leu Leu Val Ser Gly Thr Gly 1115 1120 1125	3384
cct gct gat gga aaa tac gtt cgg ttg acc gtg gac tat agc aaa Pro Ala Asp Gly Lys Tyr Val Arg Leu Thr Val Asp Tyr Ser Lys 1130 1135 1140	3429
aaa cgt tta act gtt gac tcg gtg ttt aga cag cag ctc ggt cag Lys Arg Leu Thr Val Asp Ser Val Phe Arg Gln Gln Leu Gly Gln 1145 1150 1155	3474
aga ctc ggt tcg gtt ggt ttc ttc ttg gaa aga aac ttt ggc tgt Arg Leu Gly Ser Val Gly Phe Phe Leu Glu Arg Asn Phe Gly Cys 1160 1165 1170	3519
gct caa gac gtt gaa ggt tgt ttg gtt ggt gaa gat gtt tac att Ala Gln Asp Val Glu Gly Cys Leu Val Gly Glu Asp Val Tyr Ile 1175 1180 1185	3564
gtt cag tca agg cca caa cct ctg tag Val Gln Ser Arg Pro Gln Pro Leu 1190 1195	3591

<210> 2

<211> 1196

<212> PRT

<213> Arabidopsis thaliana

<400> 2

Met Glu Ser Ile Gly Ser His Cys Cys Ser Ser Pro Phe Thr Phe Ile 1 15

Thr Arg Asn Ser Ser Ser Ser Leu Pro Arg Leu Val Asn Ile Thr His 20 25 30

Arg Val Asn Leu Ser His Gln Ser His Arg Leu Arg Asn Ser Asn Ser 35 40 45

Arg Leu Thr Cys Thr Ala Thr Ser Ser Ser Thr Ile Glu Glu Gln Arg 50 55 60

Lys Lys Lys Asp Gly Ser Gly Thr Lys Val Arg Leu Asn Val Arg Leu 65 70 75 80

Asp His Gln Val Asn Phe Gly Asp His Val Ala Met Phe Gly Ser Ala 85 90 95

Lys Glu Ile Gly Ser Trp Lys Lys Ser Pro Leu Asn Trp Ser Glu
100 105 110

Asn Gly Trp Val Cys Glu Leu Glu Leu Asp Gly Gly Gln Val Leu Glu 115 120 125

Tyr Lys Phe Val Ile Val Lys Asn Asp Gly Ser Leu Ser Trp Glu Ser 130 135 140

Gly Asp Asn Arg Val Leu Lys Val Pro Asn Ser Gly Asn Phe Ser Val 145 150 155 160

Val Cys His Trp Asp Ala Thr Arg Glu Thr Leu Asp Leu Pro Gln Glu 165 170 175

Val Gly Asn Asp Asp Asp Val Gly Asp Gly Gly His Glu Arg Asp Asn 180 185 190

His Asp Val Gly Asp Asp Arg Val Val Gly Ser Glu Asn Gly Ala Gln
195 200 205

Leu Gln Lys Ser Thr Leu Gly Gly Gln Trp Gln Gly Lys Asp Ala Ser Phe Met Arg Ser Asn Asp His Gly Asn Arg Glu Val Gly Arg Asn Trp Asp Thr Ser Gly Leu Glu Gly Thr Ala Leu Lys Met Val Glu Gly Asp Arg Asn Ser Lys Asn Trp Trp Arg Lys Leu Glu Met Val Arg Glu Val Ile Val Gly Ser Val Glu Arg Glu Glu Arg Leu Lys Ala Leu Ile Tyr Ser Ala Ile Tyr Leu Lys Trp Ile Asn Thr Gly Gln Ile Pro Cys Phe Glu Asp Gly Gly His His Arg Pro Asn Arg His Ala Glu Ile Ser Arg Leu Ile Phe Arg Glu Leu Glu His Ile Cys Ser Lys Lys Asp Ala Thr Pro Glu Glu Val Leu Val Ala Arg Lys Ile His Pro Cys Leu Pro Ser Phe Lys Ala Glu Phe Thr Ala Ala Val Pro Leu Thr Arg Ile Arg Asp Ile Ala His Arg Asn Asp Ile Pro His Asp Leu Lys Gln Glu Ile Lys His Thr Ile Gln Asn Lys Leu His Arg Asn Ala Gly Pro Glu Asp Leu Ile Ala Thr Glu Ala Met Leu Gln Arg Ile Thr Glu Thr Pro Gly Lys Tyr Ser Gly Asp Phe Val Glu Gln Phe Lys Ile Phe His Asn Glu Leu Lys Asp Phe Phe Asn Ala Gly Ser Leu Thr Glu Gln Leu Asp Ser Met Lys Ile Ser Met Asp Asp Arg Gly Leu Ser Ala Leu Asn Leu Phe Phe

Glu Cys Lys Lys Arg Leu Asp Thr Ser Gly Glu Ser Ser Asn Val Leu Glu Leu Ile Lys Thr Met His Ser Leu Ala Ser Leu Arg Glu Thr Ile Ile Lys Glu Leu Asn Ser Gly Leu Arg Asn Asp Ala Pro Asp Thr Ala Ile Ala Met Arg Gln Lys Trp Arg Leu Cys Glu Ile Gly Leu Glu Asp Tyr Phe Phe Val Leu Leu Ser Arg Phe Leu Asn Ala Leu Glu Thr Met Gly Gly Ala Asp Gln Leu Ala Lys Asp Val Gly Ser Arg Asn Val Ala Ser Trp Asn Asp Pro Leu Asp Ala Leu Val Leu Gly Val His Gln Val Gly Leu Ser Gly Trp Lys Gln Glu Glu Cys Leu Ala Ile Gly Asn Glu Leu Leu Ala Trp Arg Glu Arg Asp Leu Leu Glu Lys Glu Gly Glu Glu Asp Gly Lys Thr Ile Trp Ala Met Arg Leu Lys Ala Thr Leu Asp Arg Ala Arg Arg Leu Thr Ala Glu Tyr Ser Asp Leu Leu Gln Ile Phe Pro Pro Asn Val Glu Ile Leu Gly Lys Ala Leu Gly Ile Pro Glu Asn Ser Val Lys Thr Tyr Thr Glu Ala Glu Ile Arg Ala Gly Ile Ile Phe Gln Ile Ser Lys Leu Cys Thr Val Leu Leu Lys Ala Val Arg Asn Ser Leu Gly Ser Glu Gly Trp Asp Val Val Pro Gly Ser Thr Ser Gly Thr Leu Val Gln Val Glu Ser Ile Val Pro Gly Ser Leu Pro Ala Thr

Ser Gly Gly Pro Ile Ile Leu Leu Val Asn Lys Ala Asp Gly Asp Glu Glu Val Ser Ala Ala Asn Gly Asn Ile Ala Gly Val Met Leu Leu Gln Glu Leu Pro His Leu Ser His Leu Gly Val Arg Ala Arg Gln Glu Lys Ile Val Phe Val Thr Cys Asp Asp Asp Asp Lys Val Ala Asp Ile Arg Arg Leu Val Gly Lys Phe Val Arg Leu Glu Ala Ser Pro Ser His Val Asn Leu Ile Leu Ser Thr Glu Gly Arg Ser Arg Thr Ser Lys Ser Ser Ala Thr Lys Lys Thr Asp Lys Asn Ser Leu Ser Lys Lys Lys Thr Asp Lys Lys Ser Leu Ser Ile Asp Asp Glu Glu Ser Lys Pro Gly Ser Ser Ser Ser Asn Ser Leu Leu Tyr Ser Ser Lys Asp Ile Pro Ser Gly Gly Ile Ile Ala Leu Ala Asp Ala Asp Val Pro Thr Ser Gly Ser Lys Ser 870 · Ala Ala Cys Gly Leu Leu Ala Ser Leu Ala Glu Ala Ser Ser Lys Val His Ser Glu His Gly Val Pro Ala Ser Phe Lys Val Pro Thr Gly Val Val Ile Pro Phe Gly Ser Met Glu Leu Ala Leu Lys Gln Asn Asn Ser Glu Glu Lys Phe Ala Ser Leu Leu Glu Lys Leu Glu Thr Ala Arg Pro Glu Gly Gly Glu Leu Asp Asp Ile Cys Asp Gln Ile His Glu Val Met Lys Thr Leu Gln Val Pro Lys Glu Thr Ile Asn Ser Ile Ser Lys Ala

Phe Leu Lys Asp Ala Arg Leu Ile Val Arg Ser Ser Ala Asn Val Glu 980 985 990

Asp Leu Ala Gly Met Ser Ala Ala Gly Leu Tyr Glu Ser Ile Pro Asn 995 1000 1005

Val Ser Pro Ser Asp Pro Leu Val Phe Ser Asp Ser Val Cys Gln
1010 1020

Val Trp Ala Ser Leu Tyr Thr Arg Arg Ala Val Leu Ser Arg Arg 1025 1030 1035

Ala Ala Gly Val Ser Gln Arg Glu Ala Ser Met Ala Val Leu Val 1040 1045 1050

Gln Glu Met Leu Ser Pro Asp Leu Ser Phe Val Leu His Thr Val 1055 1060 1065

Ser Pro Ala Asp Pro Asp Ser Asn Leu Val Glu Ala Glu Ile Ala 1070 1080

Pro Gly Leu Gly Glu Thr Leu Ala Ser Gly Thr Arg Gly Thr Pro 1085 1090 1095

Trp Arg Leu Ala Ser Gly Lys Leu Asp Gly Ile Val Gln Thr Leu 1100 1110

Ala Phe Ala Asn Phe Ser Glu Glu Leu Leu Val Ser Gly Thr Gly 1115 1120 1125

Pro Ala Asp Gly Lys Tyr Val Arg Leu Thr Val Asp Tyr Ser Lys 1130 1140

Lys Arg Leu Thr Val Asp Ser Val Phe Arg Gln Gln Leu Gly Gln 1145 1150

Arg Leu Gly Ser Val Gly Phe Phe Leu Glu Arg Asn Phe Gly Cys 1160 1170

Ala Gln Asp Val Glu Gly Cys Leu Val Gly Glu Asp Val Tyr Ile 1175 1180 1185

Val Gln Ser Arg Pro Gln Pro Leu 1190 1195

<210> 3

<211> 3644

<212> DNA

<213> Oryza sativa

<220>

<221> CDS

<222> (13)..(3633)

<223>

<pre><400> 3 cgaggaggat ca atg acg tcg ctg cgg ccc ctc gaa acc tcg ctc tcc ata</pre>														51		
-9-:	,, <u>,</u>	,	Me 1	et Th	ir Se	er Le	eu Ar 5	g Pr	O Le	u G	lu Th	ir Se 10	er Le	eu Se	er Ile	71
ggc Gly	ggc Gly 15	agg Arg	ccg Pro	cgc Arg	cgt Arg	ggt Gly 20	ctc Leu	gtc Val	ctc Leu	ccg Pro	ccg Pro 25	ccc Pro	gga Gly	gtc Val	ggt Gly	99
gcg Ala 30	ggt Gly	gtg Val	ctg Leu	ctc Leu	cgc Arg 35	cgg Arg	gga Gly	gcg Ala	atg Met	gcg Ala 40	ctc Leu	cct Pro	ggg Gly	cgg Arg	cgc Arg 45	147
ggc Gly	ttc Phe	gcg Ala	tgc Cys	cgc Arg 50	ggg Gly	aga Arg	tcc Ser	gcg Ala	gcc Ala 55	tcg Ser	gcg Ala	gca Ala	gag Glu	_	aca Thr	195
aag Lys	gag Glu	aaa Lys	aag Lys 65	aga Arg	aga Arg	gat Asp	tct Ser	tca Ser 70	aag Lys	cag Gln	cca Pro	ttg Leu	gtg Val 75	cat His		243
cag Gln	gtt Val	tgt Cys 80	cta Leu	gag Glu	cac His	cag Gln	gtt Val 85	aag Lys	ttt Phe	ggt Gly	gag Glu	cat His 90	gta Val	ggc Gly	att Ile	291
atc Ile	ggt Gly 95	tcc Ser	aca Thr	aag Lys	gag Glu	ctt Leu 100	ggt Gly	tca Ser	tgg Trp	gag Glu	gag Glu 105	cag Gln	gtt Val	gaa Glu	ctg Leu	339
gaa Glu 110	tgg Trp	act Thr	aca Thr	aat Asn	ggt Gly 115	tgg Trp	gtc Val	tgc Cys	cag Gln	ctt Leu 120	aag Lys	ctc Leu	cct Pro	gga Gly	gaa Glu 125	387
aca Thr	ctt Leu	gtg Val	gag Glu	ttt Phe 130	aaa Lys	ttt Phe	gtt Val	ata Ile	ttt Phe 135	ttg Leu	gtg Val	gga Gly	gga Gly	aaa Lys 140	gat Asp	435
aaa Lys	ata Ile	tgg Trp	gaa Glu 145	gat Asp	ggt Gly	aat Asn	aac Asn	cgt Arg 150	gtt Val	gtt Val	gag Glu	ctg Leu	ccg Pro 155	aag Lys	gat Asp	483
ggt Gly	aag Lys	ttt Phe 160	Asp	Ile	Val	tgc Cys	His	Trp	Asn	Arg	Thr	gaa Glu 170	gag Glu	cca Pro	tta Leu	531
gaa Glu	ctt Leu	tta Leu	gga Gly	aca Thr	cca Pro	aag Lys	ttt Phe	gag Glu	Leu	gtc Val age	Gly	gaa Glu	gct Ala	gaa Glu	aag Lys	579

aat Asn 190	act Thr	ggc Gly	gag Glu	gat Asp	gct Ala 195	tca Ser	gca Ala	tct Ser	gta Val	act Thr 200	ttt Phe	gca Ala	cct Pro	gaa Glu	aaa Lys 205	627	
gtt Val	caa Gln	gat Asp	att Ile	tca Ser 210	gtt Val	gtt Val	gag Glu	aat Asn	ggt Gly 215	gat Asp	cca Pro	gca Ala	cca Pro	gag Glu 220	gcc Ala	675	
gag Glu	tca Ser	agc Ser	aaa Lys 225	ttt Phe	ggt Gly	ggg Gly	caa Gln	tgg Trp 230	caa Gln	gga Gly	agt Ser	aaa Lys	act Thr 235	gtt Val	ttc Phe	723	
atg Met	aga Arg	tca Ser 240	aat Asn	gag Glu	cat His	ctg Leu	aat Asn 245	aag Lys	gag Glu	gct Ala	gat Asp	agg Arg 250	atg Met	tgg Trp	gat Asp	771	
aca Thr	act Thr 255	ggg Gly	ctt Leu	gat Asp	gga Gly	ata Ile 260	gca Ala	ctg Leu	aaa Lys	ctg Leu	gtg Val 265	gag Glu	ggc Gly	gat Asp	aaa Lys	819	
gca Ala 270	tcc Ser	agg Arg	aac Asn	tgg Trp	tgg Trp 275	cgg Arg	aag Lys	tta Leu	gag Glu	gtt Val 280	gtt Val	cgc Arg	ggg Gly	ata Ile	ttg Leu 285	867	
tca Ser	gaa Glu	tct Ser	ttt Phe	gat Asp 290	gac Asp	cag Gln	agt Ser	cgt Arg	ctg Leu 295	9 9 9 Gly	gcc Ala	ctt Leu	gta Val	tac Tyr 300	tca Ser	915	
gct Ala	att Ile	tat Tyr	ctg Leu 305	aag Lys	tgg Trp	att Ile	tat Tyr	aca Thr 310	ggt Gly	cag Gln	ata Ile	tcg Ser	tgc Cys 315	ttt Phe	gaa Glu	963	
gat Asp	ggt Gly	ggc Gly 320	cac His	cat His	cgg Arg	cct Pro	aac Asn 325	aaa Lys	cat His	gct Ala	gag Glu	ata Ile 330	tcg Ser	agg Arg	caa Gln	1011	
ata Ile	ttc Phe 335	cgt Arg	gaa Glu	ctt Leu	gaa Glu	atg Met 340	atg Met	tat Tyr	tat Tyr	ggg Gly	aaa Lys 345	acc Thr	aca Thr	tca Ser	gcc Ala	1059	
aag Lys 350	gat Asp	gtt Val	ctc Leu	gtg Val	att Ile 355	cgc Arg	aaa Lys	att Ile	cat His	ccc Pro 360	ttt Phe	tta Leu	cct Pro	tca Ser	ttt Phe 365	1107	
aag Lys	tca Ser	gag Glu	ttt Phe	aca Thr 370	gcc Ala	tct Ser	gtc Val	cct Pro	cta Leu 375	aca Thr	cga Arg	att Ile	cgt Arg	gat Asp 380	att Ile	1155	
gct Ala	cac His	cgg Arg	aat Asn 385	gac Asp	atc Ile	cca Pro	cat His	gat Asp 390	ctc Leu	aag Lys	caa Gln	gaa Glu	atc Ile 395	aag Lys	cat His	1203	
act Thr	ata Ile	caa Gln 400	aac Asn	aaa Lys	ctt Leu	cat His	cgt Arg 405	aat Asn	gct Ala	gga Gly	cct Pro	gag Glu 410	gat Asp	ctt Leu	att Ile	1251	
gct Ala	aca Thr 415	gaa Glu	gtc Val	atg Met	ctt Leu	gct Ala 420	agg Arg	att Ile	act Thr	aag Lys	acc Thr 425	cct Pro	gga Gly	gaa Glu	tac Tyr	1299	
agt Ser	gaa Glu	aca Thr	ttt Phe	gtt Val	gaa Glu	caa Gln	ttc Phe	acg Thr	Ile	ttt Phe age	Tyr	agc Ser	gaa Glu	cta Leu	aaa Lys	1347	

430					435					440					445	
gat Asp	ttc Phe	ttc Phe	aat Asn	gct Ala 450	ggc Gly	agc Ser	cta Leu	ttt Phe	gag Glu 455	caa Gln	ctg Leu	gag Glu	tcc Ser	atc Ile 460	aag Lys	1395
gaa Glu	tct Ser	ctg Leu	aac Asn 465	gag Glu	tca Ser	ggc Gly	tta Leu	gaa Glu 470	gtt Val	ctc Leu	tca Ser	tcc Ser	ttt Phe 475	gtg Val	gaa Glu	1443
acc Thr	aaa Lys	agg Arg 480	agt Ser	ttg Leu	gac Asp	caa Gln	gtg Val 485	gat Asp	cat His	gca Ala	gaa Glu	gat Asp 490	ttg Leu	gat Asp	aaa Lys	1491
aat Asn	gat Asp 495	acc Thr	att Ile	caa Gln	att Ile	ttg Leu 500	atg Met	act Thr	acc Thr	ttg Leu	caa Gln 505	tca Ser	tta Leu	tct Ser	tct Ser	1539
cta Leu 510	aga Arg	tcg Ser	gtt Val	cta Leu	atg Met 515	aag Lys	ggc Gly	ctt Leu	gaa Glu	agt Ser 520	ggc Gly	ctt Leu	aga Arg	aat Asn	gat Asp 525	1587
gcg Ala	cct Pro	gat Asp	aat Asn	gct Ala 530	ata Ile	gca Ala	atg Met	cga Arg	caa Gln 535	aag Lys	tgg Trp	cgc Arg	ctt Leu	tgt Cys 540	gaa Glu	1635
att Ile	agt Ser	ctt Leu	gag Glu 545	gat Asp	tat Tyr	tca Ser	ttt Phe	gtt Val 550	ctg Leu	tta Leu	agc Ser	aga Arg	ttc Phe 555	atc Ile	aat Asn	1683
act Thr	ctt Leu	gaa Glu 560	gcc Ala	tta Leu	ggt Gly	gga Gly	tca Ser 565	gct Ala	tca Ser	ctt Leu	gca Ala	aag Lys 570	gat Asp	gta Val	gct Ala	1731
aga Arg	aat Asn 575	act Thr	act Thr	cta Leu	tgg Trp	gat Asp 580	act Thr	act Thr	ctt Leu	gat Asp	gcc Ala 585	ctt Leu	gtc Val	att Ile	ggc Gly	1779
atc Ile 590	aat Asn	caa Gln	gtt Val	agc Ser	ttt Phe 595	tca Ser	ggt Gly	tgg Trp	aaa Lys	aca Thr 600	gat Asp	gaa Glu	tgt Cys	att Ile	gcc Ala 605	1827
ata Ile	9 9 9 Gly	aat Asn	gag Glu	att Ile 610	ctt Leu	tcc Ser	tgg Trp	aag Lys	caa Gln 615	aaa Lys	ggt Gly	cta Leu	tct Ser	gaa Glu 620	agt Ser	1875
gaa Glu	ggt Gly	tgt Cys	gaa Glu 625	gat Asp	ggg Gly	aaa Lys	tat Tyr	att Ile 630	tgg Trp	tca Ser	cta Leu	aga Arg	ctt Leu 635	aaa Lys	gct Ala	1923
aca Thr	ctg Leu	gac Asp 640	aga Arg	gca Ala	cgg Arg	aga Arg	tta Leu 645	acg Thr	gaa Glu	gag Glu	tac Tyr	tct Ser 650	gaa Glu	gca Ala	ctt Leu	1971
ctt Leu	tct Ser 655	ata Ile	ttc Phe	cct Pro	gaa Glu	aaa Lys 660	gta Val	atg Met	gtt Val	att Ile	ggg Gly 665	aaa Lys	gcc Ala	ctt Leu	gga Gly	2019
ata Ile 670	cca Pro	gat Asp	aac Asn	agt Ser	gtg Val 675	aga Arg	act Thr	tac Tyr	aca Thr	gag Glu 680	gca Ala	gaa Glu	att Ile	cgt Arg	gct Ala 685	2067
ggc Gly	att Ile	gtt Val	ttt Phe	cag Gln	gta Val	tct Ser	aaa Lys	cta Leu	Cys	aca Thr age	Val	ctt Leu	cag Gln	aaa Lys	gca Ala	2115

				690					695					700		
att Ile	cga Arg	gaa Glu	gta Val 705	ctt Leu	gga Gly	tca Ser	act Thr	ggc Gly 710	tgg Trp	gat Asp	gtt Val	ctt Leu	gtt Val 715	cct Pro	gga Gly	2163
gtg Val	gcc Ala	cat His 720	gga Gly	act Thr	ctg Leu	atg Met	cgg Arg 725	gtg Val	gaa Glu	aga Arg	att Ile	ctt Leu 730	cct Pro	gga Gly	tca Ser	2211
tta Leu	cct Pro 735	tca Ser	tct Ser	gtc Val	aaa Lys	gaa Glu 740	cct Pro	gtg Val	gtt Val	cta L eu	att Ile 745	gta Val	gat Asp	aag Lys	gct Ala	2259
gat Asp 750	gga Gly	gat Asp	gaa Glu	gag Glu	gtc Val 755	aaa Lys	gct Ala	gct Ala	999 Gly	gat Asp 760	aat Asn	ata Ile	gtt Val	ggt Gly	gtt Val 765	2307
att Ile	ctt Leu	ctt Leu	cag Gln	gaa Glu 770	cta Leu	cct Pro	cac His	ctt Leu	tca Ser 775	cat His	ctt Leu	ggt Gly	gtt Val	aga Arg 780	gct Ala	2355
cgt Arg	caa Gln	gag Glu	aat Asn 785	gtt Val	gta Val	ttt Phe	gta Val	act Thr 790	tgt Cys	gaa Glu	tat Tyr	gat Asp	gac Asp 795	aca Thr	gtt Val	2403
aca Thr	gat Asp	gtg Val 800	tat Tyr	ttg Leu	ctt Leu	gag Glu	gga Gly 805	aaa Lys	tat Tyr	atc Ile	aga Arg	tta Leu 810	gaa Glu	gca Ala	tca Ser	2451
tcc Ser	atc Ile 815	aat Asn	gtc Val	aat Asn	ctc Leu	tca Ser 820	ata Ile	gtt Val	tca Ser	gaa Glu	aaa Lys 825	aat Asn	gac Asp	aat Asn	gct Ala	2499
gtc Val 830	tct Ser	aca Thr	gaa Glu	cca Pro	aat Asn 835	agt Ser	aca Thr	ggg Gly	aat Asn	cca Pro 840	ttt Phe	caa Gln	cag Gln	aaa Lys	ctc Leu 845	2547
caa Gln	aat Asn	gaa Glu	ttc Phe	tct Ser 850	cta Leu	cca Pro	tcg Ser	gat Asp	atc Ile 855	gag Glu	atg Met	cca Pro	ctg Leu	caa Gln 860	atg Met	2595
tct Ser																2643
gag Glu	ctt Leu	tca Ser 880	gaa Glu	gct Ala	tca Ser	gtg Val	gaa Glu 885	tca Ser	gct Ala	ggt Gly	gca Ala	aaa Lys 890	gct Ala	gct Ala	gca Ala	2691
tgc Cys	aga Arg 895	act Thr	ctt Leu	tct Ser	gtt Val	ctt Leu 900	gct Ala	tca Ser	ttg Leu	tct Ser	aat Asn 905	aaa Lys	gtc Val	tat Tyr	agt Ser	2739
gat Asp 910																2787
cca Pro	ttt Phe	gga Gly	tca Ser	atg Met 930	gag Glu	gat Asp	gcg Ala	ctc Leu	aag Lys 935	aaa Lys	agt Ser	gga Gly	tca Ser	ctg Leu 940	gaa Glu	2835
tcc Ser	ttt Phe	aca Thr	agc Ser	ctt Leu	cta Leu	gaa Glu	aag Lys	att Ile	Ğlu	aca Thr age	Ala	aaa Lys	gtc Val	gaa Glu	aat Asn	2883

945 950 955

ggt gaa gtt gat Gly Glu Val Asp 960	agc ctg gcg ttg gag Ser Leu Ala Leu Glu 965	cta caa gca ata att tca cat Leu Gln Ala Ile Ile Ser His 970	2931
ctt tcc cca ccg Leu Ser Pro Pro 975	gag gag act att ata Glu Glu Thr Ile Ile 980	ttt ctc aaa aga atc ttc cca Phe Leu Lys Arg Ile Phe Pro 985	2979
cag gat gtc cgg Gln Asp Val Arg 990	ttg att gtt aga tct Leu Ile Val Arg Ser 995	agt gct aat gtg gag gat ttg Ser Ala Asn Val Glu Asp Leu 1000 1005	3027
gct ggt atg tca Ala Gly Met Ser	gct gct ggt ctc tat Ala Ala Gly Leu Tyr 1010	gat tca att ccc aat gtc Asp Ser Ile Pro Asn Val 1015 1020	3072
		gct gcg gtt ggg aag gtt Ala Ala Val Gly Lys Val 1030 1035	3117
tgg gct tct tta Trp Ala Ser Leu	tac aca agg aga gcc Tyr Thr Arg Arg Ala 1040	atc cta agc cgt cga gcc Ile Leu Ser Arg Arg Ala 1045 1050	3162
gct ggt gtt tat Ala Gly Val Tyr	cag aga gac gcg aca Gln Arg Asp Ala Thr 1055	atg gct gtt ctt gtc caa Met Ala Val Leu Val Gln 1060 1065	3207
gaa ata ctg cag Glu Ile Leu Gln	cca gat ctc tcc ttc Pro Asp Leu Ser Phe 1070	gtg ctt cat act gtt tgc Val Leu His Thr Val Cys 1075 1080	3252
ccc gct gac cat Pro Ala Asp His	gac ccc aag gtt gtc Asp Pro Lys Val Val 1085	cag gct gag gtc gcc cct Gln Ala Glu Val Ala Pro 1090 1095	3297
ggg ctg ggt gaa Gly Leu Gly Glu	acg ctt gct tca gga Thr Leu Ala Ser Gly 1100	acc cgt ggc acc ccg tgg Thr Arg Gly Thr Pro Trp 1105 1110	3342
agg ctg tca tgt Arg Leu Ser Cys	aac aaa ttc gat gga Asn Lys Phe Asp Gly 1115	aaa gtt gcc act ctt gcc Lys Val Ala Thr Leu Ala 1120 1125	3387
ttt tca aat ttc Phe Ser Asn Phe	agt gag gag atg gtg Ser Glu Glu Met Val 1130	gtg cac aac tct ggt cct Val His Asn Ser Gly Pro 1135 1140	3432
gcc aat gga gaa Ala Asn Gly Glu	gta att cgt ctt act Val Ile Arg Leu Thr 1145	gtt gat tac agc aag aag Val Asp Tyr Ser Lys Lys 1150 1155	3477
cca ttg tcg gtt Pro Leu Ser Val	gat aca acc ttt agg Asp Thr Thr Phe Arg 1160	aag cag ttt ggt cag cga Lys Gln Phe Gly Gln Arg 1165 1170	3522
		cag aag ttc ggg agt gca Gln Lys Phe Gly Ser Ala 1180 1185	3567
cag gat gtg gaa Gln Asp Val Glu	ggt tgc ctg gtt ggg Gly Cys Leu Val Gly	aaa gat att ttt ata gtg Lys Asp Ile Phe Ile Val Page 16	3612

1200

3644

cca tag aagccgaatt c caa agc agg cca cag Gln Ser Arg Pro Gln Pro 1205

<210>

<211> 1206

<212> **PRT**

<213> Oryza sativa

<400> 4

Met Thr Ser Leu Arg Pro Leu Glu Thr Ser Leu Ser Ile Gly Gly Arg

Pro Arg Arg Gly Leu Val Leu Pro Pro Pro Gly Val Gly Ala Gly Val 20

Leu Leu Arg Arg Gly Ala Met Ala Leu Pro Gly Arg Arg Gly Phe Ala

Cys Arg Gly Arg Ser Ala Ala Ser Ala Ala Glu Arg Thr Lys Glu Lys 50

Lys Arg Arg Asp Ser Ser Lys Gln Pro Leu Val His Leu Gln Val Cys 70 75

Leu Glu His Gln Val Lys Phe Gly Glu His Val Gly Ile Ile Gly Ser

Thr Lys Glu Leu Gly Ser Trp Glu Glu Gln Val Glu Leu Glu Trp Thr 100 105

Thr Asn Gly Trp Val Cys Gln Leu Lys Leu Pro Gly Glu Thr Leu Val

Glu Phe Lys Phe Val Ile Phe Leu Val Gly Gly Lys Asp Lys Ile Trp 130

Glu Asp Gly Asn Asn Arg Val Val Glu Leu Pro Lys Asp Gly Lys Phe 155 145 160 150

Asp Ile Val Cys His Trp Asn Arg Thr Glu Glu Pro Leu Glu Leu Leu 175 165 170

Gly Thr Pro Lys Phe Glu Leu Val Gly Glu Ala Glu Lys Asn Thr Gly Page 17

180 185 190

Glu Asp Ala Ser Ala Ser Val Thr Phe Ala Pro Glu Lys Val Gln Asp Ile Ser Val Val Glu Asn Gly Asp Pro Ala Pro Glu Ala Glu Ser Ser Lys Phe Gly Gly Gln Trp Gln Gly Ser Lys Thr Val Phe Met Arg Ser Asn Glu His Leu Asn Lys Glu Ala Asp Arg Met Trp Asp Thr Thr Gly Leu Asp Gly Ile Ala Leu Lys Leu Val Glu Gly Asp Lys Ala Ser Arg Asn Trp Trp Arg Lys Leu Glu Val Val Arg Gly Ile Leu Ser Glu Ser Phe Asp Asp Gln Ser Arg Leu Gly Ala Leu Val Tyr Ser Ala Ile Tyr Leu Lys Trp Ile Tyr Thr Gly Gln Ile Ser Cys Phe Glu Asp Gly Gly His His Arg Pro Asn Lys His Ala Glu Ile Ser Arg Gln Ile Phe Arg Glu Leu Glu Met Met Tyr Tyr Gly Lys Thr Thr Ser Ala Lys Asp Val Leu Val Ile Arg Lys Ile His Pro Phe Leu Pro Ser Phe Lys Ser Glu Phe Thr Ala Ser Val Pro Leu Thr Arg Ile Arg Asp Ile Ala His Arg Asn Asp Ile Pro His Asp Leu Lys Gln Glu Ile Lys His Thr Ile Gln Asn Lys Leu His Arg Asn Ala Gly Pro Glu Asp Leu Ile Ala Thr Glu Val Met Leu Ala Arg Ile Thr Lys Thr Pro Gly Glu Tyr Ser Glu Thr Phe Val Glu Gln Phe Thr Ile Phe Tyr Ser Glu Leu Lys Asp Phe Phe

Page 18

Asn Ala Gly Ser Leu Phe Glu Gln Leu Glu Ser Ile Lys Glu Ser Leu Asn Glu Ser Gly Leu Glu Val Leu Ser Ser Phe Val Glu Thr Lys Arg Ser Leu Asp Gln Val Asp His Ala Glu Asp Leu Asp Lys Asn Asp Thr Ile Gln Ile Leu Met Thr Thr Leu Gln Ser Leu Ser Ser Leu Arg Ser Val Leu Met Lys Gly Leu Glu Ser Gly Leu Arg Asn Asp Ala Pro Asp Asn Ala Ile Ala Met Arg Gln Lys Trp Arg Leu Cys Glu Ile Ser Leu Glu Asp Tyr Ser Phe Val Leu Leu Ser Arg Phe Ile Asn Thr Leu Glu Ala Leu Gly Gly Ser Ala Ser Leu Ala Lys Asp Val Ala Arg Asn Thr Thr Leu Trp Asp Thr Thr Leu Asp Ala Leu Val Ile Gly Ile Asn Gln Val Ser Phe Ser Gly Trp Lys Thr Asp Glu Cys Ile Ala Ile Gly Asn Glu Ile Leu Ser Trp Lys Gln Lys Gly Leu Ser Glu Ser Glu Gly Cys Glu Asp Gly Lys Tyr Ile Trp Ser Leu Arg Leu Lys Ala Thr Leu Asp Arg Ala Arg Arg Leu Thr Glu Glu Tyr Ser Glu Ala Leu Leu Ser Ile Phe Pro Glu Lys Val Met Val Ile Gly Lys Ala Leu Gly Ile Pro Asp Asn Ser Val Arg Thr Tyr Thr Glu Ala Glu Ile Arg Ala Gly Ile Val Phe Gln Val Ser Lys Leu Cys Thr Val Leu Gln Lys Ala Ile Arg Glu

Page 19

690 695 700

Val Leu Gly Ser Thr Gly Trp Asp Val Leu Val Pro Gly Val Ala His Gly Thr Leu Met Arg Val Glu Arg Ile Leu Pro Gly Ser Leu Pro Ser Ser Val Lys Glu Pro Val Val Leu Ile Val Asp Lys Ala Asp Gly Asp Glu Glu Val Lys Ala Ala Gly Asp Asn Ile Val Gly Val Ile Leu Leu Gln Glu Leu Pro His Leu Ser His Leu Gly Val Arg Ala Arg Gln Glu Asn Val Val Phe Val Thr Cys Glu Tyr Asp Asp Thr Val Thr Asp Val Tyr Leu Leu Glu Gly Lys Tyr Ile Arg Leu Glu Ala Ser Ser Ile Asn Val Asn Leu Ser Ile Val Ser Glu Lys Asn Asp Asn Ala Val Ser Thr Glu Pro Asn Ser Thr Gly Asn Pro Phe Gln Gln Lys Leu Gln Asn Glu Phe Ser Leu Pro Ser Asp Ile Glu Met Pro Leu Gln Met Ser Lys Gln Lys Ser Lys Ser Gly Val Asn Gly Ser Phe Ala Ala Leu Glu Leu Ser Glu Ala Ser Val Glu Ser Ala Gly Ala Lys Ala Ala Ala Cys Arg Thr Leu Ser Val Leu Ala Ser Leu Ser Asn Lys Val Tyr Ser Asp Gln Gly Val Pro Ala Ala Phe Arg Val Pro Ser Gly Ala Val Ile Pro Phe Gly Ser Met Glu Asp Ala Leu Lys Lys Ser Gly Ser Leu Glu Ser Phe Thr Ser Leu Leu Glu Lys Ile Glu Thr Ala Lys Val Glu Asn Gly Glu Val Page 20

Asp Ser Leu Ala Leu Glu Leu Gln Ala Ile Ile Ser His Leu Ser Pro 965 970 975

Pro Glu Glu Thr Ile Ile Phe Leu Lys Arg Ile Phe Pro Gln Asp Val 980 985 990

Arg Leu Ile Val Arg Ser Ser Ala Asn Val Glu Asp Leu Ala Gly Met 995 1000 1005

Ser Ala Ala Gly Leu Tyr Asp Ser Ile Pro Asn Val Ser Leu Met 1010 1015 1020

Asp Pro Cys Ala Phe Gly Ala Ala Val Gly Lys Val Trp Ala Ser 1025 1030 1035

Leu Tyr Thr Arg Arg Ala Ile Leu Ser Arg Arg Ala Ala Gly Val 1040 1045 1050

Tyr Gln Arg Asp Ala Thr Met Ala Val Leu Val Gln Glu Ile Leu 1055 1060 1065

Gln Pro Asp Leu Ser Phe Val Leu His Thr Val Cys Pro Ala Asp 1070 1075 1080

His Asp Pro Lys Val Val Gln Ala Glu Val Ala Pro Gly Leu Gly 1085 1090 1095

Glu Thr Leu Ala Ser Gly Thr Arg Gly Thr Pro Trp Arg Leu Ser 1100 1110

Cys Asn Lys Phe Asp Gly Lys Val Ala Thr Leu Ala Phe Ser Asn 1115 1120 1125

Phe Ser Glu Glu Met Val Val His Asn Ser Gly Pro Ala Asn Gly 1130 1140

Glu Val Ile Arg Leu Thr Val Asp Tyr Ser Lys Lys Pro Leu Ser 1145 1150 1155

Val Asp Thr Thr Phe Arg Lys Gln Phe Gly Gln Arg Leu Ala Ala 1160 1170

Ile Gly Gln Tyr Leu Glu Gln Lys Phe Gly Ser Ala Gln Asp Val 1175 1180 1185

Glu Gly Cys Leu Val Gly Lys Asp Ile Phe Ile Val Gln Ser Arg Page 21 1190 1195 1200

Pro Gln Pro 1205

<210> 5

<211> 12

<212> PRT

<213> Oryza sativa, Arabidopsis thaliana, Sorghum bicolor

<400> 5

Leu Pro His Leu Ser His Leu Gly Val Arg Ala Arg
1 10

<210> 6

<211> 7

<212> PRT

<213> Hordeum vulgare

<400> 6

Ser Arg Arg Val Ala Gly Val 5

<210> 7

<211> 7

<212> PRT

<213> Hordeum vulgare

<400> 7

Val Glu Ala Glu Val Ala Pro 1 5

<210> 8

<211> 9

<212> PRT

<213> Hordeum vulgare

His Thr Val Ser Pro Ser Asp His Asp <210> 9 <211> 807 <212> DNA <213> Hordeum vulgare <220> <221> CDS (3)..(590)<222> <223> <400> 9 47 cg gca cga gga gtc ctc ccc aat gtg agc ctc tcg gac cca acc aac Ala Arg Gly Val Leu Pro Asn Val Ser Leu Ser Asp Pro Thr Asn 15 10 95 ttc ggg tct gca gta gcg cgg gtc tgg gcc tcg ctg tac act cgg agg Phe Gly Ser Ala Val Ala Arg Val Trp Ala Ser Leu Tyr Thr Arg Arg 20 30 143 gcc atc ctc agc cgc cgg gtg gct ggc gtg ccc cag agg gac gcc aag Ala Ile Leu Ser Arg Arg Val Ala Gly Val Pro Gln Arg Asp Ala Lys atg gct gtc ctg gtg cag gag atg ctg gag cca gag cta tcc ttc gtg 191 Met Ala Val Leu Val Gli Gli Met Leu Glu Pro Gli Leu Ser Phe Val 55 50 60 ctc cac acg gtc agc ccc tcg gac cac gac acc agg gtc gtc gag gct 239 Leu His Thr Val Ser Pro Ser Asp His Asp Thr Arg Val Val Glu Ala 65 287 gag gtt gcc ccg ggg ctg ggc gag acc ctt gcc gct ggc acc cgc ggc Glu Val Āla Pro Gly Leu Gly Glu Thr Leu Āla Āla Gly Thr Arg Gly 80 90 85 335 acc ccg tgg cgt ctc tcc tgc gac aag ttc gac acc gac gtc gcc acc Thr Pro Trp Arg Leu Ser Cys Asp Lys Phe Asp Thr Asp Val Ala Thr 110 100 383 ctg gcc ttc gcc aac ttc agt gag gag atg cgg gtg ctc ggc tcg ggc Leu Ala Phe Ala Asn Phe Ser Glu Glu Met Arg Val Leu Gly Ser Gly 115 120 ccc gcc gac ggc gag gtg gtg agg ctc act gtc gac tac agc acg aag 431 Pro Ala Asp Gly Glu Val Val Arg Leu Thr Val Asp Tyr Ser Thr Lys 135 130 140

<400> 8

ctg cte Leu Lei 14	u Ser														479
gcc gcc Ala Ala 160	c gtg a Val	ggg Gly	cag Gln	tac Tyr 165	ctg Leu	gag Glu	cag Gln	agg Arg	ttc Phe 170	999 Gly	agc Ser	gcc Ala	cag Gln	gac Asp 175	527
gtg gag Val Gl	g ggc u Gly	tgc Cys	atg Met 180	gtc val	tgg Trp	gaa Glu	gac Asp	atc Ile 185	tac Tyr	ata Ile	gtg Val	cag Gln	agc Ser 190	atg Met	575
cca ca Pro Gl			tag	agto	catco	cgt a	aataa	atgti	tt ag	gatga	agcaa	a agi	tttt	ggtt	630
ggtgaa	ataa a	aatti	tgcc	ga aa	aatco	ccat	g gca	aaaa	taag	tcag	ggta	tga a	agago	cccgcc	690
tgcgaa	acca a	actga	attci	ta aa	ataat	tgtti	t tga	aatto	cgtg	ttta	aaat	tat (ggga	cgtgaa	750
caatga	tttc (cttg	gaat	gc at	tgcat	ttgta	a ag	tttt	aaaa	aaaa	aaaa	aaa a	aaaa	aaa	807
<210>	10														
<211>	195														
<212>	PRT														
<213>	Hord	eum '	vulga	are											

<400> 10

Ala Arg Gly Val Leu Pro Asn Val Ser Leu Ser Asp Pro Thr Asn Phe 10 15

Gly Ser Ala Val Ala Arg Val Trp Ala Ser Leu Tyr Thr Arg Arg Ala 20 25 30

Ile Leu Ser Arg Arg Val Ala Gly Val Pro Gln Arg Asp Ala Lys Met 35 40 45

Ala Val Leu Val Gln Glu Met Leu Glu Pro Glu Leu Ser Phe Val Leu 50 60

His Thr Val Ser Pro Ser Asp His Asp Thr Arg Val Val Glu Ala Glu 65 70 75 80

Val Ala Pro Gly Leu Gly Glu Thr Leu Ala Ala Gly Thr Arg Gly Thr 85 90 95

Pro Trp Arg Leu Ser Cys Asp Lys Phe Asp Thr Asp Val Ala Thr Leu 100 105 110

Ala Phe Ala Asn Phe Ser Glu Glu Met Arg Val Leu Gly Ser Gly Pro Page 24 Ala Asp Gly Glu Val Val Arg Leu Thr Val Asp Tyr Ser Thr Lys Leu 130 135 140

125

Leu Ser Val Asp Arg Thr Phe Arg Gln Lys Phe Gly Gln Arg Leu Ala 145 150 150

Ala Val Gly Gln Tyr Leu Glu Gln Arg Phe Gly Ser Ala Gln Asp Val 165 170 175

Glu Gly Cys Met Val Trp Glu Asp Ile Tyr Ile Val Gln Ser Met Pro 180 185 190

Gln Pro Leu 195

<210> 11

<211> 9

<212> PRT

<213> Solanum tuberosum

<400> 11

Pro Glu Glu Cys Lys Ala Val Gly Asn 5

<210> 12

<211> 7

<212> PRT

<213> Solanum tuberosum

<400> 12

Thr Glu Glu Tyr Ser Glu Thr

<210> 13

<211> 7

<212> PRT

<213> Solanum tuberosum

```
<400> 13
Arg Phe Val Asn Ala Val Glu
1 5
<210> 14
<211>
<212>
      PRT
<213> Solanum tuberosum
<400> 14
Glu Gly Ser Glu Asp Gly Lys
1
<210> 15
<211> 403
<212>
       DNA
<213> Solanum tuberosum
<220>
<221>
      CDS
<222> (1)..(402)
<223>
<400> 15
                                                                       48
gcg gat gct tca ata gct atg cgt cag aag tgg cgt ctc tgc gaa atc
Ala Asp Ala Ser Ile Ala Met Arg Gln Lys Trp Arg Leu Cys Glu Ile
                                                         15
                                                                       96
ggg ctt gaa gac tat gca ttt gtt ctt ttg agc agg ttt gtg aat gca
Gly Leu Glu Asp Tyr Ala Phe Val Leu Leu Ser Arg Phe Val Asn Ala
                                                     30
            20
                                 25
                                                                       144
gtt gaa gct cta ggc gga gct gat tgg ctt gca gag aat gta aca gtg
Val Glu Ala Leu Gly Gly Ala Asp Trp Leu Ala Glu Asn Val Thr Val
        35
                                                                       192
aaa aac att agt tct tgg aat gat cca att gga gca ctt aca gtt gga
Lys Asn Ile Ser Ser Trp Asn Asp Pro Ile Gly Ala Leu Thr Val Gly
    50
                                             60
                                                                       240
atc caa cag cta ggt ata tct ggt tgg aag ccc gag gaa tgc aaa gct
Ile Gln Gln Leu Gly Ile Ser Gly Trp Lys Pro Glu Glu Cys Lys Ala
65
                                         75
                                                             80
                    70
```

gtt gga aat gaa ctt ttg tca tgg aaa gaa agg ggt att tca gaa att Val Gly Asn Glu Leu Leu Ser Trp Lys Glu Arg Gly Ile Ser Glu Ile 85 90 95
gaa ggc agc gaa gat ggt aag act ata tgg gca tta aga cta aaa gcg Glu Gly Ser Glu Asp Gly Lys Thr Ile Trp Ala Leu Arg Leu Lys Ala 100 105 110
act ctt gat aga agt cga agg tta act gag gag tat tcc gag aca ctt Thr Leu Asp Arg Ser Arg Arg Leu Thr Glu Glu Tyr Ser Glu Thr Leu 115 120 125
ctc caa ata ttc cct gaa a Leu Gln Ile Phe Pro Glu 130
<210> 16
<211> 134
<212> PRT
<213> Solanum tuberosum
<400> 16
Ala Asp Ala Ser Ile Ala Met Arg Gln Lys Trp Arg Leu Cys Glu Ile 1 5 10 15
Gly Leu Glu Asp Tyr Ala Phe Val Leu Leu Ser Arg Phe Val Asn Ala 20 25 30
Val Glu Ala Leu Gly Gly Ala Asp Trp Leu Ala Glu Asn Val Thr Val 35 40 45
Lys Asn Ile Ser Ser Trp Asn Asp Pro Ile Gly Ala Leu Thr Val Gly 50 55 60
Ile Gln Gln Leu Gly Ile Ser Gly Trp Lys Pro Glu Glu Cys Lys Ala 65 70 75 80
Val Gly Asn Glu Leu Leu Ser Trp Lys Glu Arg Gly Ile Ser Glu Ile 85 90 95
Glu Gly Ser Glu Asp Gly Lys Thr Ile Trp Ala Leu Arg Leu Lys Ala 100 105 110
Thr Leu Asp Arg Ser Arg Arg Leu Thr Glu Glu Tyr Ser Glu Thr Leu 115 120 125
Leu Gln Ile Phe Pro Glu 130

<210> 17

<211> 7

<212> PRT

<213> Sorghum bicolor

<400> 17

Asp Gly Gly His His Arg Pro 5

<210> 18

<211> 8

<212> PRT

<213> Sorghum bicolor

<400> 18

Asp Ala Pro Asp Ser Ala Ile Ala 1

<210> 19

<211> 9

<212> PRT

<213> Sorghum bicolor

<400> 19

Ile Pro Glu Asn Ser Val Arg Thr Tyr

<210> 20

<211> 6

<212> PRT

<213> Sorghum bicolor

<400> 20

Val Asn Lys Ala Asp Gly 5

<210> 21 <211> 1526 <212> DNA <213> Sorghum bicolor <220> <221> CDS <222> (2)..(1525)<223> <400> 21 49 g cac gag gct gaa tat gtt cat gat cag agt cac ctg gag gct ctt aca His Glu Ala Glu Tyr Val His Asp Gln Ser His Leu Glu Ala Leu Thr 10 15 97 tat tct gca ata tat cta aag tgg ata tat act ggt caa ata cca tgc Tyr Ser Ala Ile Tyr Leu Lys Trp Ile Tyr Thr Gly Gln Ile Pro Cys 20 145 ttt gag gat ggt ggt cac cat cga ccc aat aaa cat gct gag ata tcc Phe Glu Asp Gly Gly His His Arg Pro Asn Lys His Ala Glu Ile Ser 193 agg caa att ttt cgt gaa att gaa agg ata tac tat ggg gaa aac aca Arg Gln Ile Phe Arg Glu Ile Glu Arg Ile Tyr Tyr Gly Glu Asn Thr 55 50 241 tca gct cag gat ttg ctt gtg ata cgc aag att cat cct tgt cta cct Ser Ala Gln Asp Leu Leu Val Ile Arg Lys Ile His Pro Cys Leu Pro 289 tca ttt aaa tca gaa ttt act gcc tct gtt cct cta aca cga att cgt Ser Phe Lys Ser Glu Phe Thr Ala Ser Val Pro Leu Thr Arg Ile Arg 90 337 gat att gct cat cgt aat gac ata cca cat gat ctc aag caa gaa atc Asp Ile Ala His Arg Asn Asp Ile Pro His Asp Leu Lys Gln Glu Ile 100 105 110 385 aag cat act ata caa aac aag ctt cac cgg aat gcc ggc cct gag gat Lys His Thr Ile Gln Asn Lys Leu His Arg Asn Ala Gly Pro Glu Asp 115 120 433 ctt att gct act gaa gcc atg ctt gct agg att act aag act cct gga Leu Ile Ala Thr Glu Ala Met Leu Ala Arg Ile Thr Lys Thr Pro Gly 130 135 140 481 gag tac agt gaa gct ttt gtt gaa caa ttc aag acg ttt tat agt gaa Glu Tyr Ser Glu Ala Phe Val Glu Gln Phe Lys Thr Phe Tyr Ser Glu 145 155 160 150 529 tta aaa gat ttc ttc aat gct ggc agc cta ctg gag caa gtg caa tcc

Leu Lys Asp Phe Phe Asn Ala Gly Ser Leu Leu Glu Gln Val Gln Ser

Page 29

				165					170					175		
atc Ile	gag Glu	caa Gln	tct Ser 180	ttg Leu	gat Asp	gag Glu	tct Ser	ggc Gly 185	tta Leu	gaa Glu	gct Ala	ctc Leu	tca Ser 190	tcc Ser	ttt Phe	577
ctg Leu	aaa Lys	acc Thr 195	aaa Lys	aag Lys	aat Asn	tta Leu	gac Asp 200	caa Gln	ctg Leu	gaa Glu	gat Asp	gca Ala 205	aaa Lys	gat Asp	ttg Leu	625
gat Asp	gaa Glu 210	aat Asn	ggt Gly	ggc Gly	gtt Val	caa Gln 215	gtt Val	ttg Leu	ttg Leu	aaa Lys	gcc Ala 220	ttg Leu	ctg Leu	tcg Ser	tta Leu	673
tct Ser 225	tat Tyr	cta Leu	aga Arg	tca Ser	att Ile 230	cta Leu	atg Met	aag Lys	ggt Gly	ctg Leu 235	gaa Glu	agt Ser	ggc Gly	ctt Leu	aga Arg 240	721
aat Asn	gat Asp	gct Ala	cca Pro	gat Asp 245	agt Ser	gct Ala	att Ile	gca Ala	atg Met 250	cga Arg	caa Gln	aag Lys	tgg Trp	cgt Arg 255	ctt Leu	769
tgt Cys	gag Glu	atc Ile	999 Gly 260	ctt Leu	gaa Glu	gat Asp	tat Tyr	tcg Ser 265	ttt Phe	gta Val	ttg Leu	tta Leu	agt Ser 270	aga Arg	tac Tyr	817
atc Ile	aat Asn	gct Ala 275	ctt Leu	gaa Glu	gct Ala	ttg Leu	ggt Gly 280	gga Gly	tca Ser	gct Ala	tca Ser	ctt Leu 285	gca Ala	gag Glu	ggt Gly	865
ctt Leu	cct Pro 290	aca Thr	aat Asn	aca Thr	agt Ser	cta Leu 295	tgg Trp	gat Asp	gat Asp	gcc Ala	ctt Leu 300	gat Asp	gcc Ala	ctt Leu	gtc Val	913
att Ile 305	ggc Gly	ata Ile	aat Asn	caa Gln	gtt Val 310	agc Ser	ttt Phe	tca Ser	gga Gly	tgg Trp 315	aaa Lys	cca Pro	aat Asn	gag Glu	tgt Cys 320	961
		ata Ile														1009
gaa Glu	ttt Phe	gaa Glu	ggc Gly 340	agt Ser	gag Glu	gat Asp	gga Gly	aag Lys 345	tat Tyr	att Ile	tgg Trp	gca Ala	ctg Leu 350	aga Arg	ctc Leu	1057
aaa Lys	gcc Ala	act Thr 355	ctt Leu	gat Asp	aga Arg	tca Ser	cga Arg 360	aga Arg	cta Leu	aca Thr	gaa Glu	gaa Glu 365	tac Tyr	tct Ser	gaa Glu	1105
		ctt Leu													gcc Ala	1153
ctt Leu 385	gga Gly	ata Ile	cca Pro	gag Glu	aac Asn 390	agt Ser	gtg Val	aga Arg	aca Thr	tac Tyr 395	act Thr	gaa Glu	gct Ala	gaa Glu	att Ile 400	1201
		ggt Gly													tta Leu	1249
		act Thr							Ser		Trp				gtt Val	1297

			420					425					430			
cct Pro	gga Gly	gtg Val 435	gcc Ala	cat His	gga Gly	gcc Ala	ttg Leu 440	ata Ile	cag Gln	gtt Val	gaa Glu	aga Arg 445	ata Ile	gct Ala	cct Pro	1345
gga Gly	tca Ser 450	ttg Leu	cca Pro	tca Ser	tcc Ser	atc Ile 455	aaa Lys	gaa Glu	cct Pro	gtc Val	gtg Val 460	cta Leu	gtt val	gta Val	aac Asn	1393
aag Lys 465	gct Ala	gat Asp	gga Gly	gat Asp	gaa Glu 470	gag Glu	gtc Val	aaa Lys	gct Ala	gct Ala 475	ggg Gly	gat Asp	aac Asn	ata Ile	gtg Val 480	1441
ggt Gly	gtt Val	att Ile	ctt Leu	cta Leu	caa Gln	gaa Glu	tta Leu	cct Pro	cac His	cta Leu	tca Ser	cat His	ctt Leu	ggt Gly	gtt Val	1489

aga gct cgt caa gag aaa gtt gta ttt gta act tgc g

Arg Ala Arg Gln Glu Lys Val Val Phe Val Thr Cys

500

505

490

495

<210> 22

<211> 508

<212> PRT

< <213> Sorghum bicolor

485

<400> 22

His Glu Ala Glu Tyr Val His Asp Gln Ser His Leu Glu Ala Leu Thr 1 10 15

Tyr Ser Ala Ile Tyr Leu Lys Trp Ile Tyr Thr Gly Gln Ile Pro Cys 20 25 30

Phe Glu Asp Gly Gly His His Arg Pro Asn Lys His Ala Glu Ile Ser 35 40 45

Arg Gln Ile Phe Arg Glu Ile Glu Arg Ile Tyr Tyr Gly Glu Asn Thr 50 55 60

Ser Ala Gln Asp Leu Leu Val Ile Arg Lys Ile His Pro Cys Leu Pro 65 75 80

Ser Phe Lys Ser Glu Phe Thr Ala Ser Val Pro Leu Thr Arg Ile Arg 85 90 95

Asp Ile Ala His Arg Asn Asp Ile Pro His Asp Leu Lys Gln Glu Ile 100 105 110

Lys His Thr Ile Gln Asn Lys Leu His Arg Asn Ala Gly Pro Glu Asp Page 31 115 120 125

Leu Ile Ala Thr Glu Ala Met Leu Ala Arg Ile Thr Lys Thr Pro Gly Glu Tyr Ser Glu Ala Phe Val Glu Gln Phe Lys Thr Phe Tyr Ser Glu Leu Lys Asp Phe Phe Asn Ala Gly Ser Leu Leu Glu Gln Val Gln Ser Ile Glu Gln Ser Leu Asp Glu Ser Gly Leu Glu Ala Leu Ser Ser Phe Leu Lys Thr Lys Lys Asn Leu Asp Gln Leu Glu Asp Ala Lys Asp Leu Asp Glu Asn Gly Gly Val Gln Val Leu Leu Lys Ala Leu Leu Ser Leu Ser Tyr Leu Arg Ser Ile Leu Met Lys Gly Leu Glu Ser Gly Leu Arg Asn Asp Ala Pro Asp Ser Ala Ile Ala Met Arg Gln Lys Trp Arg Leu Cys Glu Ile Gly Leu Glu Asp Tyr Ser Phe Val Leu Leu Ser Arg Tyr Ile Asn Ala Leu Glu Ala Leu Gly Gly Ser Ala Ser Leu Ala Glu Gly Leu Pro Thr Asn Thr Ser Leu Trp Asp Asp Ala Leu Asp Ala Leu Val Ile Gly Ile Asn Gln Val Ser Phe Ser Gly Trp Lys Pro Asn Glu Cys Thr Ala Ile Val Asn Glu Leu Leu Ser Trp Lys Gln Lys Gly Leu Ser Glu Phe Glu Gly Ser Glu Asp Gly Lys Tyr Ile Trp Ala Leu Arg Leu Lys Ala Thr Leu Asp Arg Ser Arg Arg Leu Thr Glu Glu Tyr Ser Glu Ala Leu Leu Ser Ile Phe Pro Glu Lys Val Lys Val Leu Gly Lys Ala

Page 32

Leu Gly Ile Pro Glu Asn Ser Val Arg Thr Tyr Thr Glu Ala Glu Ile 385 390 395 400

Arg Ala Gly Val Ile Phe His Val Ser Lys Leu Cys Thr Val Leu Leu 405 410 415

Lys Ala Thr Arg Ala Val Leu Gly Ser Ser Val Trp Asp Val Leu Val 420 425 430

Pro Gly Val Ala His Gly Ala Leu Ile Gln Val Glu Arg Ile Ala Pro 435 440 445

Gly Ser Leu Pro Ser Ser Ile Lys Glu Pro Val Val Leu Val Val Asn 450 455 460

Lys Ala Asp Gly Asp Glu Glu Val Lys Ala Ala Gly Asp Asn Ile Val 465 470 475 480

Gly Val Ile Leu Leu Gln Glu Leu Pro His Leu Ser His Leu Gly Val 485 490 495

Arg Ala Arg Gln Glu Lys Val Val Phe Val Thr Cys 500 505

<210> 23

<211> 8

<212> PRT

<213> Triticum aestivum

<400> 23

Arg Asn Asp Ala Thr Asp Ala Gly
1

<210> 24

<211> 8

<212> PRT

<213> Triticum aestivum

<400> 24

Gly Asn Thr Ser Val Trp Asp Asp

<212> DNA

<223>

<213> Triticum aestivum

<220>
<221> CDS
<222> (1)..(507)

<400> 25 48 aat ggc gct ttt gtc gaa caa ttt caa ata ttt tat agc gaa cta aaa Asn Gly Ala Phe Val Glu Gln Phe Gln Ile Phe Tyr Ser Glu Leu Lys 10 gac ttc ttt aat gcc ggc agc ctg ttt gaa caa ctg gaa tcc atc aag 96 Asp Phe Phe Asn Ala Gly Ser Leu Phe Glu Gln Leu Glu Ser Ile Lys 30 20 144 gaa tot ttg aat gat tot ggo tta gaa goa otg toa toa ttt gto aaa Glu Ser Leu Asn Asp Ser Gly Leu Glu Ala Leu Ser Ser Phe Val Lys 35 192 acc aaa cag agt ttg gac caa gtg gat gct gcg aac att caa gtt gtg Thr Lys Gln Ser Leu Asp Gln Val Asp Ala Ala Asn Ile Gln Val Val 240 atg aag acc ttg cag tca ttg tct tca ttg aga tca gtt cta atg aag Met Lys Thr Leu Gln Ser Leu Ser Ser Leu Arg Ser Val Leu Met Lys 65 80 288 ggc ctt gaa agt ggc ctt aga aat gat gcg act gat gcc ggt ata gca Gly Leu Glu Ser Gly Leu Arg Asn Asp Ala Thr Asp Ala Gly Ile Ala 336 atg cga caa aag tgg cgc ctt tgt gag att ggt ctt gag gat tat tct Met Arg Gln Lys Trp Arg Leu Cys Glu Ile Gly Leu Glu Asp Tyr Ser 105 100 384 ttt gtt ttg tta agc aga tat atc aat ggt ctt gaa gct tca ggt gga Phe Val Leu Leu Ser Arg Tyr Ile Asn Gly Leu Glu Ala Ser Gly Gly 115 120 tca gct tca ctt gca caa tgt gtg gct gga aat aca agt gta tgg gac 432 Ser Ala Ser Leu Ala Gln Cys Val Ala Gly Asn Thr Ser Val Trp Asp 130 135 480 gat acc ctt gat gcc ctt att att ggc gtc aat caa gtt agc ttt tca Asp Thr Leu Asp Ala Leu Ile Ile Gly Val Asn Gln Val Ser Phe Ser 145 150 155 160

ggt tgg aag cca gag gaa tgc att gct at Gly Trp Lys Pro Glu Glu Cys Ile Ala 165

<210> 26

<211> 169

<212> PRT

<213> Triticum aestivum

<400> 26

Asn Gly Ala Phe Val Glu Gln Phe Gln Ile Phe Tyr Ser Glu Leu Lys
1 10 15

Asp Phe Phe Asn Ala Gly Ser Leu Phe Glu Gln Leu Glu Ser Ile Lys 20 25 30

Glu Ser Leu Asn Asp Ser Gly Leu Glu Ala Leu Ser Ser Phe Val Lys 35 40 45

Thr Lys Gln Ser Leu Asp Gln Val Asp Ala Ala Asn Ile Gln Val Val 50 60

Met Lys Thr Leu Gln Ser Leu Ser Ser Leu Arg Ser Val Leu Met Lys 70 75 80

Gly Leu Glu Ser Gly Leu Arg Asn Asp Ala Thr Asp Ala Gly Ile Ala 85 90 95

Met Arg Gln Lys Trp Arg Leu Cys Glu Ile Gly Leu Glu Asp Tyr Ser 100 105 110

Phe Val Leu Leu Ser Arg Tyr Ile Asn Gly Leu Glu Ala Ser Gly Gly 115 120 125

Ser Ala Ser Leu Ala Gln Cys Val Ala Gly Asn Thr Ser Val Trp Asp 130 135 140

Asp Thr Leu Asp Ala Leu Ile Ile Gly Val Asn Gln Val Ser Phe Ser 145 150 155 160

Gly Trp Lys Pro Glu Glu Cys Ile Ala 165

<210> 27 <211> 27

<212> DNA

<213> Synthetic	
<400> 27 gactcaacca cataacaca aaagatc	27
<210> 28 <211> 21 <212> DNA <213> Synthetic	
<400> 28 tggtaacgag gcaaatgcag a	21
<210> 29 <211> 25 <212> DNA <213> Synthetic	
<400> 29 atctcttatc acaccacctc caatg	25
<210> 30 <211> 24 <212> DNA <213> Synthetic	
<400> 30 ggaaccgata atgcctacat gctc	24
<210> 31 <211> 39 <212> DNA <213> Synthetic	
<400> 31 aaaactcgag gaggatcaat gacgtcgctg cggcccctc	39
<210> 32 <211> 21 <212> DNA <213> Synthetic	
<400> 32 ccaggttaag tttggtgagc a	21
<210> 33 <211> 22 <212> DNA <213> Synthetic	
<400> 33 caaagcacga tatctgacct gt	22
<210> 34 <211> 22 <212> DNA <213> Synthetic	

ttgttcgcgg gatattgtca ga	22
<210> 35 <211> 24 <212> DNA <213> Synthetic	
<400> 35 gacaagggca tcaagagtag tatc	24
<210> 36 <211> 20 <212> DNA <213> Synthetic	
<400> 36 atgatgcgcc tgataatgct	20
<210> 37 <211> 21 <212> DNA <213> Synthetic	
<400> 37 ggcaaacagt atgaagcacg a	21
<210> 38 <211> 21 <212> DNA <213> Synthetic <400> 38 catttggatc aatggaggat g	21
<210> 39 <211> 23 <212> DNA <213> Synthetic	
<400> 39 ctatggctgt ggcctgcttt gca	23
<210> 40 <211> 33 <212> DNA <213> Synthetic	
<400> 40 aaaactcgag ctatggctgt ggcctgcttt gca	33
<210> 41 <211> 28 <212> DNA <213> Synthetic	

<400> 41 aaaacaattg gcgcctggag ggaggaga	28
<210> 42 <211> 35 <212> DNA <213> Synthetic	
<400> 42 aaaacaattg atgatcaatc agacaatcac tagaa	35
<210> 43 <211> 63 <212> DNA <213> Synthetic	
<pre><400> 43 tgcaggctgc agagctccta ggctcgagtt aacactagta agcttaatta agatatcat tac</pre>	t 60 63
<210> 44 <211> 67 <212> DNA <213> Synthetic	
<400> 44 aattgtaaat gatatcttaa ttaagcttac tagtgttaac tcgagcctag gagctctgc gcctgca	a 60 67
<210> 45 <211> 28 <212> DNA <213> Synthetic	
<400> 45 ttttctcgag gtccgccttg tttctcct	28
<210> 46 <211> 26 <212> DNA <213> Synthetic	
<400> 46 ttttctcgag ctgcacgggt ccagga	26